RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/830,691A
Source: IFW0
Date Processed by STIC: 1-6-4

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RE-RUN

IFWO

RAW SEQUENCE LISTING DATE: 01/06/2005
PATENT APPLICATION: US/09/830,691A TIME: 14:49:08

Input Set : N:\Crf4\Refhold\09_folder\I830691A.raw
Output Set: N:\CRF4\01062005\I830691A.raw

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      1 <110> APPLICANT: Choi, Eui-Sung
      2       Rhee, Sang-Ki
      3       Sohn, Jung-Hoon
      4       Park, Soo-Dong
      5       Lee, Yoon-Hyoung
      6       Lee, Seung-Jae
      7       Jang, Jae-Kweon
      8       Choi, Seok-Keun
      9       Son, Young-Rok
     10 <120> TITLE OF INVENTION: VECTOR FOR THE TRANSFORMATION OF PHAFFIA
     11       RHODOZYMA AND PROCESS OF TRANSFORMATION THEREBY
     12 <130> FILE REFERENCE: 118.12-US-WO
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/830,691A
     14 <141> CURRENT FILING DATE: 2001-04-26
     15 <150> PRIOR APPLICATION NUMBER: KR 1998/46547
     16 <151> PRIOR FILING DATE: 1998-10-31
     17 <150> PRIOR APPLICATION NUMBER: PCT/KR99/00265
     18 <151> PRIOR FILING DATE: 1999-05-29
     19 <160> NUMBER OF SEQ ID NOS: 20
     20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     22 <210> SEQ ID NO: 1
     23 <211> LENGTH: 1223
     24 <212> TYPE: DNA
     25 <213> ORGANISM: Phaffia rhodozyma
     26 <400> SEQUENCE: 1
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     28       caaatattcc agtgcatcga aagagtttgt ggataaacgc gacagtttca agggaaagag 120
     29       tcgatggaca gatttggaag acttagccgg tcaaggaact tggggatcac gtggcggagg 180
     30       actcatcaga agaagtcggg atttgtttga tcatagtggg atcaagacaa actggaggat 240
     31       atggctcgcc ttggaaggga atctccggcc tggattcgag gatccgaaag ttgtacgtat 300
     32       ggaaaagctt acacggcttg gatttattat ctttcatagg aacctactgc aagggtaagg 360
     33       cttgcaagaa gcacacgtaa gtcgcttatc ctctccactc tttcatggca tattgtcaac 420
     34       gactggacaa cgcgtccgtt ttgaaacaag tgacttacct gtgaaatttg attctacacc 480
     35       tgtatttagc cctcacaagg tacatatcac atcctcccac cccaccctgc ccaacttctt 540
     36       cagttcatct tgctctcggt ttccacattc cctgatgacc tccttgtatg ttctttgcga 600
     37       acgtttgttt ctgtttctgt aggtgaccca gtacaagaag ggaaaggact ccatcttcgc 660
     38       ccagggaaag cgacgatacg accgaaagca gtccggttac ggaggtcaga ccaagcccgt 720
     39       tttccacaag aaggctaaga ccaccaagaa ggtcgtcctt cgattggcgg tatttttgtt 780
     40       tattttgaat tctttttgtg tatgcagact tttgatgatt atgctcctct gtcgtttttt 840
     41       ctcttcaaac agagtgctcc gtctgcagtt cgttcttcct tccaaccaaa acttcaacta 900
     42       cagacatcat aaacagacat cttacttcgg tgttctctct ttttttccgc agagtacaag 960
     43       atgcagatga ccctcaagcg atgcaagcac ttcgagcttg gaggagacaa gaagaccaag 1020
     44       ggttcgtctt ttgtccatat attctctggt tcacttctta tgttcctaac gtacttgttt 1080
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45       cctttttggt tcggatgttg tttctatcgg tggtgttttc ttttctttgg atgcattatc 1140
46       atttatcgtg ttggactgtt ttcctctgct cgtttctttc tcctctgtac ttgtgcttct 1200
47       caggagccgc catctctttc taa                                         1223
49  <210> SEQ ID NO: 2
50  <211> LENGTH: 350
51  <212> TYPE: DNA
52  <213> ORGANISM: Phaffia rhodozyma
53  <220> FEATURE:
54  <221> NAME/KEY: CDS
55  <222> LOCATION: (30)...(347)
56  <400> SEQUENCE: 2
57       cccttcaagt ctcgtctcaa tcagtcaag atg gtc aac gtt ccc aag act cga   53
58                                     Met Val Asn Val Pro Lys Thr Arg
59                                      1               5
60       cga acc tac tgc aag ggt aag gct tgc aag aag cac acc cct cac aag   101
61       Arg Thr Tyr Cys Lys Gly Lys Ala Cys Lys Lys His Thr Pro His Lys
62            10                  15                  20
63       gtg acc cag tac aag aag gga aag gac tcc atc ttc gcc cag gga aag   149
64       Val Thr Gln Tyr Lys Lys Gly Lys Asp Ser Ile Phe Ala Gln Gly Lys
65        25                  30                  35                  40
66       cga cga tac gac cga aag cag tcc ggt tac gga ggt cag acc aag ccc   197
67       Arg Arg Tyr Asp Arg Lys Gln Ser Gly Tyr Gly Gly Gln Thr Lys Pro
68                          45                  50                  55
69       gtt ttc cac aag aag gct aag acc acc aag aag gtc gtc ctt cga ttg   245
70       Val Phe His Lys Lys Ala Lys Thr Thr Lys Lys Val Val Leu Arg Leu
71                     60                  65                  70
72       gag tgc tcc gtc tgc aag tac aag atg cag atg acc ctc aag cga tgc   293
73       Glu Cys Ser Val Cys Lys Tyr Lys Met Gln Met Thr Leu Lys Arg Cys
74                  75                  80                  85
75       aag cac ttc gag ctt gga gga gac aag aag acc aag gga gcc gcc atc   341
76       Lys His Phe Glu Leu Gly Gly Asp Lys Lys Thr Lys Gly Ala Ala Ile
77            90                  95                  100
78       tct ttc taa                                                     350
79       Ser Phe
80       105
82  <210> SEQ ID NO: 3
83  <211> LENGTH: 106
84  <212> TYPE: PRT
85  <213> ORGANISM: Phaffia rhodozyma
86  <400> SEQUENCE: 3
87       Met Val Asn Val Pro Lys Thr Arg Arg Thr Tyr Cys Lys Gly Lys Ala
88        1               5                  10                  15
89       Cys Lys Lys His Thr Pro His Lys Val Thr Gln Tyr Lys Lys Gly Lys
90                   20                  25                  30
91       Asp Ser Ile Phe Ala Gln Gly Lys Arg Arg Tyr Asp Arg Lys Gln Ser
92               35                  40                  45
93       Gly Tyr Gly Gly Gln Thr Lys Pro Val Phe His Lys Lys Ala Lys Thr
94           50                  55                  60
95       Thr Lys Lys Val Val Leu Arg Leu Glu Cys Ser Val Cys Lys Tyr Lys
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```
     96       65                  70                  75                  80
     97       Met Gln Met Thr Leu Lys Arg Cys Lys His Phe Glu Leu Gly Gly Asp
     98                       85                  90                  95
     99       Lys Lys Thr Lys Gly Ala Ala Ile Ser Phe
     100                  100                 105
     102 <210> SEQ ID NO: 4
     103 <211> LENGTH: 741
     104 <212> TYPE: DNA
     105 <213> ORGANISM: Phaffia rhodozyma
     106 <220> FEATURE:
     107 <221> NAME/KEY: misc_feature
     108 <222> LOCATION: (0)...(0)
     109 <223> OTHER INFORMATION: n=a, t, c, or g
     110 <400> SEQUENCE: 4
     111       ctcgagtgga cggtggcaat ggcattcgtg tcgttggtgc tcactcgcaa cccaagcagt 60
     112       cgcttacccg gggtagcctc cgggtgggcg cgatgatttg tggtgtggat tccttcccta 120
     113       tgggtagaac gacgcgcaac caatcattcg gagaaccgct ccgttgtagc cgaccagtct 180
     114       gattgatcaa catgccagca cgtcctccgg gacggagact ggcggggatc gtacctcatc 240
     115       tggaatcgct ggctcaatgg tagtagtctt cacgatcggc catgagggca gtctaggtgg 300
W--> 116       gttcgcctgc cgaagactgt gtgagtgtgc tganaactaa ttgagtaccg ggggataagg 360
     117       caaggcgtgt ntggttgccg gtggctgtga gcgagtttgc tgcaaagcga ttcaatgcac 420
     118       cccggcttgg ccagcgcgct gcgtcacgaa acacactaaa cggttgacgc cataaagtaa 480
     119       taacacactc aagtttgtgg tcccgggtgg gcctctgtgc ctgcgtggga cccgacggga 540
     120       gaggaaaacg ttctgtggcc ctctcctctg tggatagtta cctggttgat cctgccagta 600
     121       gtcatatgct tgtctcaaag attaagccat gcatgtctaa gtataaacaa attcatactg 660
     122       tgaaactgcg aatggctcat taaatcagtt atagtttatt tgatggtacc ttgctacatg 720
     123       gataactgtg gtaattctag a                                          741
     125 <210> SEQ ID NO: 5
     126 <211> LENGTH: 23
     127 <212> TYPE: DNA
     128 <213> ORGANISM: Artificial Sequence
     129 <220> FEATURE:
     130 <223> OTHER INFORMATION: CYH1, a PCR primer for the cloning of L41 genomic
     131       DNA fragment
     132 <220> FEATURE:
     133 <221> NAME/KEY: misc_feature
     134 <222> LOCATION: (0)...(0)
     135 <223> OTHER INFORMATION: n=a, t, c, or g
     136 <400> SEQUENCE: 5
W--> 137       cgcgtagtta aygtnccnaa rac                                        23
     139 <210> SEQ ID NO: 6
     140 <211> LENGTH: 25
     141 <212> TYPE: DNA
     142 <213> ORGANISM: Artificial Sequence
     143 <220> FEATURE:
     144 <223> OTHER INFORMATION: CYH3, a PCR primer for the cloning of L41 genomic
     145       DNA fragment
     146 <400> SEQUENCE: 6
     147       cccgggtytt ggcyttyttr tgraa                                      25
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Input Set : N:\Crf4\Refhold\09_folder\I830691A.raw
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149 <210> SEQ ID NO: 7
150 <211> LENGTH: 24
151 <212> TYPE: DNA
152 <213> ORGANISM: Artificial Sequence
153 <220> FEATURE:
154 <223> OTHER INFORMATION: 3' RACE primer
155 <400> SEQUENCE: 7
156 ggtcagacca agcaagtttt tcac 24
158 <210> SEQ ID NO: 8
159 <211> LENGTH: 24
160 <212> TYPE: DNA
161 <213> ORGANISM: Artificial Sequence
162 <220> FEATURE:
163 <223> OTHER INFORMATION: 5' RACE primer
164 <400> SEQUENCE: 8
165 gtgaaaaact tgcttggtct gacc 24
167 <210> SEQ ID NO: 9
168 <211> LENGTH: 24
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
171 <220> FEATURE:
172 <223> OTHER INFORMATION: sense primer for the mutagenesis of L41 gene
173 <400> SEQUENCE: 9
174 ggtcagacca agcaagtttt tcac 24
176 <210> SEQ ID NO: 10
177 <211> LENGTH: 24
178 <212> TYPE: DNA
179 <213> ORGANISM: Artificial Sequence
180 <220> FEATURE:
181 <223> OTHER INFORMATION: antisense primer for the mutagenesis of L41 gene
182 <400> SEQUENCE: 10
183 gtgaaaaact tgcttggtct gacc 24
185 <210> SEQ ID NO: 11
186 <211> LENGTH: 20
187 <212> TYPE: DNA
188 <213> ORGANISM: Artificial Sequence
189 <220> FEATURE:
190 <223> OTHER INFORMATION: a PCR primer corresponding to 18S rDNA
191 <400> SEQUENCE: 11
192 tcctagtaag cgcaagtcat 20
194 <210> SEQ ID NO: 12
195 <211> LENGTH: 20
196 <212> TYPE: DNA
197 <213> ORGANISM: Artificial Sequence
198 <220> FEATURE:
199 <223> OTHER INFORMATION: a PCR primer corresponding to 18S rDNA
200 <400> SEQUENCE: 12
201 ttcggccaag gaaagaaact 20
203 <210> SEQ ID NO: 13

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Input Set : N:\Crf4\Refhold\09_folder\I830691A.raw
Output Set: N:\CRF4\01062005\I830691A.raw

<211> LENGTH: 20
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: a PCR primer corresponding to 28S rDNA
<400> SEQUENCE: 13
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<210> SEQ ID NO: 14
<211> LENGTH: 20
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: a PCR primer corresponding to 28S rDNA
<400> SEQUENCE: 14
gctataacac atccggagat 20
<210> SEQ ID NO: 15
<211> LENGTH: 2192
<212> TYPE: DNA
<213> ORGANISM: Phaffia rhodozyma
<400> SEQUENCE: 15
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atctcaattt ttcaaatact tgccaacact ttcatatatt cacaccaaaa aaagtcagat 180
tggcccacaa agtcagatac acgctcgatc gtcgacgggt tcaagcactt tgtcaggcga 240
aagaaaggcc acagcaccac ccttcaagtc tcgtctcaat caggttcgtc tagcttttg 300
tgtgcaagga tttaccgtct tgatggattt gttcgttgaa agagaggaaa gaacatgctg 360
aactgacgaa agtgtgaaca aaaaattgtg atttttтcat tgtgtttcgc tggtctcctt 420
gctgggttgg gttggatcgg atttatcttc tgtgttggat ggaaaaccct gaatgttctt 480
ttcttggaca tcttctaaac tcgacaaaac gattcattcc tccgtactgc tctggttctg 540
ccttttgaa tcgcatcgat aaattcttcc ctcggaacgt tcgatcaatc tccgtcaaac 600
ttatcatcca aaaatctctt ctcgactgcc gccttgctcc ttttcttcgt tcttttcctta 660
atccgctttc gactaccctc cttctcttca cactcatagt caagatggtc aacgttccca 720
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tcgaaagagt ttgtggataa acgcgacagt ttcaagggaa agagtcgatg gacagatttg 840
gaagacttag ccggtcaagg aacttgggga tcacgtggcg gaggactcat cagaagaagt 900
cgggatttgt ttgatcatag tgggatcaag acaaactgga ggatatggct cgccttggaa 960
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cgttttgaaa caagtgactt acctgtgaaa tttgattcta cacctgtatt tagccctcac 1200
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tacgaccgaa agcagtccgg ttacggaggt cagaccaagc ccgtttttca caagaaggct 1440
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tgtgtatgca gacttttgat gattatgctc ctctgtcgtt ttttctcttc aaacagagtg 1560
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acatcttact tcggtgttct ctctttttтt ccgcagagta caagatgcag atgaccctca 1680
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 01/06/2005
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; N Pos. 334,371
Seq#:5; N Pos. 15,18

VERIFICATION SUMMARY DATE: 01/06/2005
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Input Set : N:\Crf4\Refhold\09_folder\I830691A.raw
Output Set: N:\CRF4\01062005\I830691A.raw

L:13 M:270 C: Current Application Number differs, Wrong Format
L:116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:300
M:341 Repeated in SeqNo=4
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0